



2

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/517,941
Source: Pt 110
Date Processed by STIC: 12/28/04

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.2 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<<http://www.uspto.gov/ebc/efs/downloads/documents.htm>> , EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry, Federal Express, United Parcel Service, or other delivery service (EFFECTIVE 06/05/04):
U.S. Patent and Trademark Office, 220 20th Street S., Customer Window, Mail Stop Sequence, Crystal Plaza Two, Lobby, Room 1B03, Arlington, VA 22202

Raw Sequence Listing Error Summary

ERROR DETECTED

SUGGESTED CORRECTION

SERIAL NUMBER:

10/517,941

ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE

- 1 Wrapped Nucleics
 Wrapped Aminos The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."

- 2 Invalid Line Length The rules require that a line not exceed 72 characters in length. This includes white spaces.

- 3 Misaligned Amino
 Numbering The numbering under each 5th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.

- 4 Non-ASCII The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.

- 5 Variable Length Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.

- 6 PatentIn 2.0
 "bug" A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.

- 7 Skipped Sequences
 (OLD RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence:
 (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading)
 (xi) SEQUENCE DESCRIPTION: SEQ ID NO:X: (insert SEQ ID NO where "X" is shown)
 This sequence is intentionally skipped

 Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.

- 8 Skipped Sequences
 (NEW RULES) Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence.
 <210> sequence id number
 <400> sequence id number
 000

- 9 Use of n's or Xaa's
 (NEW RULES) Use of n's and/or Xaa's have been detected in the Sequence Listing.
 Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present.
 In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.

- 10 Invalid <213>
 Response Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence

- 11 Use of <220> Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses.
 Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section.
 (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)

- 12 PatentIn 2.0
 "bug" Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.

- 13 Misuse of n/Xaa "n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid



PCT

RAW SEQUENCE LISTING

DATE: 12/28/2004

PATENT APPLICATION: US/10/517,941

TIME: 11:51:26

Input Set : D:\0077 (OX40).ST25.txt

Output Set: N:\CRF4\12282004\J517941.raw

3 <110> APPLICANT: Crucell Holland B.V.
 4 Bakker, Alexander B.H.
 5 Meester-Rood, Pauline M.L.
 6 Bakker, Adrianus Q.
 8 <120> TITLE OF INVENTION: AGONISTIC BINDING MOLECULES TO THE HUMAN OX40 RECEPTOR
 10 <130> FILE REFERENCE: 0077 WO 00 ORD
 C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/517,941
 C--> 12 <141> CURRENT FILING DATE: 2004-12-13
 12 <150> PRIOR APPLICATION NUMBER: PCT/NL02/00389
 13 <151> PRIOR FILING DATE: 2002-06-13
 15 <160> NUMBER OF SEQ ID NOS: 54
 17 <170> SOFTWARE: PatentIn version 3.1
 19 <210> SEQ ID NO: 1
 20 <211> LENGTH: 769
 21 <212> TYPE: DNA
 22 <213> ORGANISM: Artificial sequence
 24 <220> FEATURE:
 25 <223> OTHER INFORMATION: (scFv SC02008) what is the source of the
 27 <220> FEATURE: genetic material?
 28 <221> NAME/KEY: CDS (see item 11 on Euro summary
 29 <222> LOCATION: (3)..(767) sheet)
 30 <223> OTHER INFORMATION:
 W--> 33 <400> 1
 34 cc atg gct gag gtg cag ctg gtg gag tct ggg gga ggc ttg gtc cag 47
 35 Met Ala Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln
 36 1 5 10 15
 38 cct gga ggg tcc ctg aga ctc tcc tgt gca gcc tct gga ttc acc ttt 95
 39 Pro Gly Gly Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe
 40 20 25 30
 42 agc aac tac acg atg aac tgg gtc cgc cag gcg ccc ggg aag ggg ctg 143
 43 Ser Asn Tyr Thr Met Asn Trp Val Arg Gln Ala Pro Gly Lys Gly Leu
 44 35 40 45
 46 gag tgg gtc tca gct att agt ggt agt ggt ggt agc aca tac tac gca 191
 47 Glu Trp Val Ser Ala Ile Ser Gly Ser Gly Gly Ser Thr Tyr Tyr Ala
 48 50 55 60
 50 gac tcc gtg aag ggc cgg ttc acc atc tcc aga gac aat tcc aag aac 239
 51 Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn
 52 65 70 75
 54 acg ctg tat ctg caa atg aac agc ctg aga gcc gag gac acg gcc gtg 287
 55 Thr Leu Tyr Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val
 56 80 85 90 95
 58 tat tac tgt gcc aaa gac cgc tac tcc cag gtg cac tac gcg ttg gat 335
 59 Tyr Tyr Cys Ala Lys Asp Arg Tyr Ser Gln Val His Tyr Ala Leu Asp

pp 1-5
 Does Not Comply
 Corrected Diskette Needed

RAW SEQUENCE LISTING

DATE: 12/28/2004

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TIME: 11:51:26

Input Set : D:\0077 (OX40).ST25.txt

Output Set: N:\CRF4\12282004\J517941.raw

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60          100          105          110
62 tac tgg ggc cag ggc acc ctg gtg acc gtg ctc gag ggt acc gga ggt      383
63 Tyr Trp Gly Gln Gly Thr Leu Val Thr Val Leu Glu Gly Thr Gly Gly
64          115          120          125
66 tcc ggc gga acc ggg tct ggg act ggt acg agc gag ctc gac atc cag      431
67 Ser Gly Gly Thr Gly Ser Gly Thr Gly Thr Ser Glu Leu Asp Ile Gln
68          130          135          140
70 atg acg cag tct cca gac tca ctg ccc gtc acc cct gga gag ccg gcc      479
71 Met Thr Gln Ser Pro Asp Ser Leu Pro Val Thr Pro Gly Glu Pro Ala
72          145          150          155
74 tcc atc tcc tgc agg tct agt cag agc ctc ctg cat agt aat gga tac      527
75 Ser Ile Ser Cys Arg Ser Ser Gln Ser Leu Leu His Ser Asn Gly Tyr
76 160          165          170          175
78 aac tat ttg gat tgg tac ctg cag aag gca ggg cag tct cca cag ctc      575
79 Asn Tyr Leu Asp Trp Tyr Leu Gln Lys Ala Gly Gln Ser Pro Gln Leu
80          180          185          190
82 ctg atc tat ttg ggt tct aat cgg gcc tcc ggg gtc cct gac agg ttc      623
83 Leu Ile Tyr Leu Gly Ser Asn Arg Ala Ser Gly Val Pro Asp Arg Phe
84          195          200          205
86 agt ggc agt gga tca ggc aca gat ttt aca ctg aaa atc agc aga gtg      671
87 Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Lys Ile Ser Arg Val
88          210          215          220
90 gag gct gag gat gtt ggg gtt tat tac tgc cag cag tac tac aac cac      719
91 Glu Ala Glu Asp Val Gly Val Tyr Tyr Cys Gln Gln Tyr Tyr Asn His
92          225          230          235
94 ccg acg acc ttc ggc cag ggc acc aaa ctg gaa atc aaa cgc gcg gcc      767
95 Pro Thr Thr Phe Gly Gln Gly Thr Lys Leu Glu Ile Lys Arg Ala Ala
96 240          245          250          255
98 gc      769

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101 <210> SEQ ID NO: 2

102 <211> LENGTH: 255

103 <212> TYPE: PRT

104 <213> ORGANISM: Artificial sequence

106 <220> FEATURE:

107 <223> OTHER INFORMATION: scFv SC02008

109 <400> SEQUENCE: 2

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111 Met Ala Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln Pro
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115 Gly Gly Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser
116          20          25          30
119 Asn Tyr Thr Met Asn Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu
120          35          40          45
123 Trp Val Ser Ala Ile Ser Gly Ser Gly Gly Ser Thr Tyr Tyr Ala Asp
124          50          55          60
127 Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn Thr
128 65          70          75          80
131 Leu Tyr Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala Val Tyr
132          85          90          95
135 Tyr Cys Ala Lys Asp Arg Tyr Ser Gln Val His Tyr Ala Leu Asp Tyr

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RAW SEQUENCE LISTING

DATE: 12/28/2004

PATENT APPLICATION: US/10/517,941

TIME: 11:51:26

Input Set : D:\0077 (OX40).ST25.txt

Output Set: N:\CRF4\12282004\J517941.raw

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136          100          105          110
139 Trp Gly Gln Gly Thr Leu Val Thr Val Leu Glu Gly Thr Gly Gly Ser
140          115          120          125
143 Gly Gly Thr Gly Ser Gly Thr Gly Thr Ser Glu Leu Asp Ile Gln Met
144          130          135          140
147 Thr Gln Ser Pro Asp Ser Leu Pro Val Thr Pro Gly Glu Pro Ala Ser
148 145          150          155          160
151 Ile Ser Cys Arg Ser Ser Gln Ser Leu Leu His Ser Asn Gly Tyr Asn
152          165          170          175
155 Tyr Leu Asp Trp Tyr Leu Gln Lys Ala Gly Gln Ser Pro Gln Leu Leu
156          180          185          190
159 Ile Tyr Leu Gly Ser Asn Arg Ala Ser Gly Val Pro Asp Arg Phe Ser
160          195          200          205
163 Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Lys Ile Ser Arg Val Glu
164          210          215          220
167 Ala Glu Asp Val Gly Val Tyr Tyr Cys Gln Gln Tyr Tyr Asn His Pro
168 225          230          235          240
171 Thr Thr Phe Gly Gln Gly Thr Lys Leu Glu Ile Lys Arg Ala Ala
172          245          250          255
175 <210> SEQ ID NO: 3
176 <211> LENGTH: 775
177 <212> TYPE: DNA
178 <213> ORGANISM: Artificial sequence
180 <220> FEATURE:
181 <223> OTHER INFORMATION: scFv SC02009
183 <220> FEATURE:
184 <221> NAME/KEY: CDS
185 <222> LOCATION: (3)..(773)
186 <223> OTHER INFORMATION:

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191 Met Ala Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln
192 1          5          10          15
194 cct ggg ggg tcc ctg aga ctc tcc tgt gca gcc tct gga ttc acc ttc      95
195 Pro Gly Gly Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe
196          20          25          30
198 agc ggc tac tct atg aac tgg gtc cgc cag gcg ccc ggg aag ggg ctg      143
199 Ser Gly Tyr Ser Met Asn Trp Val Arg Gln Ala Pro Gly Lys Gly Leu
200          35          40          45
202 gag tgg gtt ggc cgt act aga aac aaa gct aac agt tac acc aca gaa      191
203 Glu Trp Val Gly Arg Thr Arg Asn Lys Ala Asn Ser Tyr Thr Thr Glu
204          50          55          60
206 tac gcc gcg tct gtg aaa ggc aga ttc acc atc tca aga gat gat tca      239
207 Tyr Ala Ala Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asp Ser
208          65          70          75
210 aag aac tca ctg tat ctg caa atg aac agt ctg aga gcc gag gac aca      287
211 Lys Asn Ser Leu Tyr Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr
212 80          85          90          95
214 gcc gtg tat tac tgt gcc aaa gac cgc tac gtc aac acg tcg aac gcg      335

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DATE: 12/28/2004

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TIME: 11:51:26

Input Set : D:\0077 (OX40).ST25.txt

Output Set: N:\CRF4\12282004\J517941.raw

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215 Ala Val Tyr Tyr Cys Ala Lys Asp Arg Tyr Val Asn Thr Ser Asn Ala
216          100          105          110
218 ttc gat tac tgg ggc cag ggc acc ctg gtg acc gtg ctc gag ggt acc      383
219 Phe Asp Tyr Trp Gly Gln Gly Thr Leu Val Thr Val Leu Glu Gly Thr
220          115          120          125
222 gga ggt tcc ggc gga acc ggg tct ggg act ggt acg agc gag ctc gac      431
223 Gly Gly Ser Gly Gly Thr Gly Ser Gly Thr Gly Thr Ser Glu Leu Asp
224          130          135          140
226 atc cag atg aca cag tct cca gac tca ctg ccc gtc acc cct gga gag      479
227 Ile Gln Met Thr Gln Ser Pro Asp Ser Leu Pro Val Thr Pro Gly Glu
228          145          150          155
230 ccg gcc tcc atc tcc tgc aga tct agt cag agc ctc ctg cat agt aat      527
231 Pro Ala Ser Ile Ser Cys Arg Ser Ser Gln Ser Leu Leu His Ser Asn
232 160          165          170          175
234 gga tac aac tat ttg gat tgg tac ctg cag aag cca ggg cag tct cca      575
235 Gly Tyr Asn Tyr Leu Asp Trp Tyr Leu Gln Lys Pro Gly Gln Ser Pro
236          180          185          190
238 cag ctc ctg atc tat ttg ggt tct aat cgg gcc tcc ggg gtc cct gac      623
239 Gln Leu Leu Ile Tyr Leu Gly Ser Asn Arg Ala Ser Gly Val Pro Asp
240          195          200          205
242 agg ttc agt ggc agt gga tca ggc aca gat ttt aca ctg aaa atc agc      671
243 Arg Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Lys Ile Ser
244          210          215          220
246 aga gtg gag gct cac cat gtt ggg gtt tat tac tgc cag cag tac ccg      719
247 Arg Val Glu Ala His His Val Gly Val Tyr Tyr Cys Gln Gln Tyr Pro
248          225          230          235
250 ctg ggc ccg ccc acc ttc ggc cag ggc acc aaa ctg gaa atc aaa cgc      767
251 Leu Gly Pro Pro Thr Phe Gly Gln Gly Thr Lys Leu Glu Ile Lys Arg
252 240          245          250          255
254 gcg gcc gc      775
255 Ala Ala
259 <210> SEQ ID NO: 4
260 <211> LENGTH: 257
261 <212> TYPE: PRT
262 <213> ORGANISM: Artificial sequence
264 <220> FEATURE:
265 <223> OTHER INFORMATION: scFv SC02009
267 <400> SEQUENCE: 4
269 Met Ala Glu Val Gln Leu Val Glu Ser Gly Gly Gly Leu Val Gln Pro
270 1          5          10          15
273 Gly Gly Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Ser
274          20          25          30
277 Gly Tyr Ser Met Asn Trp Val Arg Gln Ala Pro Gly Lys Gly Leu Glu
278          35          40          45
281 Trp Val Gly Arg Thr Arg Asn Lys Ala Asn Ser Tyr Thr Thr Glu Tyr
282          50          55          60
285 Ala Ala Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asp Ser Lys
286 65          70          75          80
289 Asn Ser Leu Tyr Leu Gln Met Asn Ser Leu Arg Ala Glu Asp Thr Ala

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DATE: 12/28/2004

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TIME: 11:51:26

Input Set : D:\0077 (OX40).ST25.txt

Output Set: N:\CRF4\12282004\J517941.raw

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290          85          90          95
293 Val Tyr Tyr Cys Ala Lys Asp Arg Tyr Val Asn Thr Ser Asn Ala Phe
294          100          105          110
297 Asp Tyr Trp Gly Gln Gly Thr Leu Val Thr Val Leu Glu Gly Thr Gly
298          115          120          125
301 Gly Ser Gly Gly Thr Gly Ser Gly Thr Gly Thr Ser Glu Leu Asp Ile
302          130          135          140
305 Gln Met Thr Gln Ser Pro Asp Ser Leu Pro Val Thr Pro Gly Glu Pro
306 145          150          155          160
309 Ala Ser Ile Ser Cys Arg Ser Ser Gln Ser Leu Leu His Ser Asn Gly
310          165          170          175
313 Tyr Asn Tyr Leu Asp Trp Tyr Leu Gln Lys Pro Gly Gln Ser Pro Gln
314          180          185          190
317 Leu Leu Ile Tyr Leu Gly Ser Asn Arg Ala Ser Gly Val Pro Asp Arg
318          195          200          205
321 Phe Ser Gly Ser Gly Ser Gly Thr Asp Phe Thr Leu Lys Ile Ser Arg
322          210          215          220
325 Val Glu Ala His His Val Gly Val Tyr Tyr Cys Gln Gln Tyr Pro Leu
326 225          230          235          240
329 Gly Pro Pro Thr Phe Gly Gln Gly Thr Lys Leu Glu Ile Lys Arg Ala
330          245          250          255
333 Ala

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337 <210> SEQ ID NO: 5

338 <211> LENGTH: 736

339 <212> TYPE: DNA

340 <213> ORGANISM: Artificial sequence

342 <220> FEATURE:

343 <223> OTHER INFORMATION: scFv SC02010

345 <220> FEATURE:

346 <221> NAME/KEY: CDS

347 <222> LOCATION: (3)..(734)

348 <223> OTHER INFORMATION:

W--> 351 <400> 5

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352 cc atg gct gag gtg cag ctg gtg gag tct ggg gga ggc ttg atc cag      47
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354 1          5          10          15
356 cct ggg ggg tcc ctg aga ctc tcc tgt gca gcc tct gga ttc acc ttc      95
357 Pro Gly Gly Ser Leu Arg Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe
358          20          25          30
360 agc ggc tac cct atg aac tgg gtc cgc cag gcg ccc ggg aag ggg ctg      143
361 Ser Gly Tyr Pro Met Asn Trp Val Arg Gln Ala Pro Gly Lys Gly Leu
362          35          40          45
364 gag tgg gtg gca gtt ata tca tat gat gga agt aat aaa tac tac gca      191
365 Glu Trp Val Ala Val Ile Ser Tyr Asp Gly Ser Asn Lys Tyr Tyr Ala
366          50          55          60
368 gac tcc gtg aag ggc cga ttc acc atc tcc aga gac aat tcc aag aac      239
369 Asp Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asn Ser Lys Asn
370          65          70          75
372 acg ctg tat ctg caa atg aac agc ctg aga gct gag gac aca gcc gtg      287

```

IMPORTANT

The types of errors shown exist throughout the Sequence Listing. Please check subsequent sequences for similar errors.

VERIFICATION SUMMARY

DATE: 12/28/2004

PATENT APPLICATION: US/10/517,941

TIME: 11:51:27

Input Set : D:\0077 (OX40).ST25.txt

Output Set: N:\CRF4\12282004\J517941.raw

L:12 M:270 C: Current Application Number differs, Replaced Current Application No
L:12 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:33 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:1,Line#:30
L:189 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:3,Line#:186
L:351 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:5,Line#:348
L:505 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:7,Line#:502
L:659 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:9,Line#:656
L:813 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:11,Line#:810
L:969 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:13,Line#:966
L:1123 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:15,Line#:1120
L:2157 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:33,Line#:2154
L:2199 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:35,Line#:2196
L:2241 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:37,Line#:2238